12

SEQUENCE CORRECTION REPORT
PATENT APPLICATION US/07/978,891B

LINE	ORIGINAL TEXT	CORRECTED TEXT
4	(1)GENERAL INFORMATION	(1) GENERAL INFORMATION:
12	(iv)CORRESPONDING ADDRESS:	(iv) CORRESPONDENCE ADDRESS:
349	(3)INFORMATION FOR SEQ ID NO: 2:	(2) INFORMATION FOR SEQ ID NO: 2:
676	(4)INFORMATION FOR SEQ ID NO: 3:	(2) INFORMATION FOR SEQ ID NO: 3:
695	(5)INFORMATION FOR SEQ ID NO: 4:	(2) INFORMATION FOR SEQ ID NO: 4:
714	(6)INFORMATION FOR SEQ ID NO: 5:	(2) INFORMATION FOR SEQ ID NO: 5:
749	(7)INFORMATION FOR SEQ ID NO: 6:	(2) INFORMATION FOR SEQ ID NO: 6:
768	(8)INFORMATION FOR SEQ ID NO: 7:	(2) INFORMATION FOR SEQ ID NO: 7:
787	(9)INFORMATION FOR SEQ ID NO: 8:	(2) INFORMATION FOR SEQ ID NO: 8:

MANDATORY IDENTIFIER THAT WAS NOT FOUND

PRIOR APPLICATION DATA
APPLICATION NUMBER
FILING DATE

LINE ERROR   ORIGINAL TEXT

OK

LINE	ERROR	ORIGINAL TEXT
30	Wrong application Serial Number	(A)APPLICATION NUMBER: 07/978,891
691	Entered and Calc. Seq. Length differ	(ix)SEQUENCE DESCRIPTION: SEQ ID NO: 3
729	Entered and Calc. Seq. Length differ	(ix)SEQUENCE DESCRIPTION: SEQ ID NO: 5
764	Entered and Calc. Seq. Length differ	(ix)SEQUENCE DESCRIPTION: SEQ ID NO: 6
783	Entered and Calc. Seq. Length differ	(ix)SEQUENCE DESCRIPTION: SEQ ID NO: 7
802	Entered and Calc. Seq. Length differ	(ix)SEQUENCE DESCRIPTION: SEQ ID NO: 8

Au 1805

Patent Application US/07/978,891B

PP. 14 ⟶ 16

1	SEQUENCE LISTING
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4	(1)GENERAL INFORMATION
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6	(i)APPLICANT: Darrell Anderson, Nabil Hanna, John Leonard, Roland Newman and Mitchell R
7	
8	(ii)TITLE OF INVENTION: THERAPEUTIC APPLICATION OF CHIMERIC ANTIBODY TO HUMAN B LYMPHOCY
9	
10	(iii)NUMBER OF SEQUENCES: 8
11	
12	(iv)CORRESPONDING ADDRESS:
13	
14	(A)ADDRESSEE: IDEC Pharmaceuticals Corporation
15	(B)STREET:11099 N. Torrey Pines Road, #160
16	(C)CITY:La Jolla
17	(D)STATE:California
18	(E)COUNTRY:USA
19	(F)ZIP:92037
20	
21	(v)COMPUTER READABLE FORM:
22	
23	(A)MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb
24	(B)COMPUTER: Macintosh
25	(C)OPERATING SYSTEM: MS.DOS
26	(D)SOFTWARE: Microsoft Word 5.0
27	
28	(vi)CURRENT APPLICATION DATA:
29	
30	(A)APPLICATION NUMBER: 07/978,891
31	(B)FILING DATE: 13 NOV 1992
32	(C)CLASSIFICATION: 424
33	
34	(viii)ATTORNEY/AGENT INFORMATION:
35	
36	(A)NAME: Burgoon, Richard P. Jr.
37	(B)REGISTRATION NUMBER: 34,787
38	(C)REFERENCE/DOCKET NUMBER:
39	
40	(ix)TELECOMMUNICATION INFORMATION:
41	
42	(A)TELEPHONE: (619) 458-0600
43	(B)TELEFAX: (619) 546-9274
44	
45	(2)INFORMATION FOR SEQ ID NO: 1:
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47	(i)SEQUENCE CHARACTERISTICS:
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49	(A)LENGTH: 8540 bases
50	(B)TYPE: nucleic acid
51	(C)STRANDEDNESS: single
52	(D)TOPOLOGY: circular

53
54   (ii)MOLECULE TYPE: DNA (genomic)
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56   (iii)HYPOTHETICAL: no
57
58   (iv)ANTI-SENSE: no
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60   (ix)SEQUENCE DESCRIPTION: SEQ ID NO: 1:
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Raw Sequence Listing

Patent Application US/07/978,891B

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419   AGCCTCAGCA GCACCCTGAC GCTGAGCAAA GCAGACTACG AGAAACACAA AGTCTACGCC   1620
420
421   TGCGAAGTCA CCCATCAGGG CCTGAGCTCG CCCGTCACAA AGAGCTTCAA CAGGGGAGAG   1680
422
423   TGTTGAATTC AGATCCGTTA ACGGTTACCA ACTACCTAGA CTGGATTCGT GACAACATGC   1740
424
425   GGCCGTGATA TCTACGTATG ATCAGCCTCG ACTGTGCCTT CTAGTTGCCA GCCATCTGTT   1800
426
427   GTTTGCCCCT CCCCCGTGCC TTCCTTGACC CTGGAAGGTG CCACTCCCAC TGTCCTTTCC   1860
428
429   TAATAAAATG AGGAAATTGC ATCGCATTGT CTGAGTAGGT GTCATTCTAT TCTGGGGGGT   1920
430
431   GGGGTGGGGC AGGACAGCAA GGGGGAGGAT TGGGAAGACA ATAGCAGGCA TGCTGGGGAT   1980
432
433   GCGGTGGGCT CTATGGAACC AGCTGGGGCT CGACAGCTAT GCCAAGTACG CCCCCTATTG   2040
434
435   ACGTCAATGA CGGTAAATGG CCCGCCTGGC ATTATGCCCA GTACATGACC TTATGGGACT   2100
436
437   TTCCTACTTG GCAGTACATC TACGTATTAG TCATCGCTAT TACCATGGTG ATGCGGTTTT   2160
438
439   GGCAGTACAT CAATGGGCGT GGATAGCGGT TTGACTCACG GGGATTTCCA AGTCTCCACC   2220
440
441   CCATTGACGT CAATGGGAGT TTGTTTTGGC ACCAAAATCA ACGGGACTTT CCAAAATGTC   2280
442
443   GTAACAACTC CGCCCCATTG ACGCAAATGG GCGGTAGGCG TGTACGGTGG GAGGTCTATA   2340
444
445   TAAGCAGAGC TGGGTACGTC CTCACATTCA GTGATCAGCA CTGAACACAG ACCCGTCGAC   2400
446
447   ATGGGTTGGA GCCTCATCTT GCTCTTCCTT GTCGCTGTTG CTACGCGTGT CCTGTCCCAG   2460
448
449   GTACAACTGC AGCAGCCTGG GGCTGAGCTG GTGAAGCCTG GGCCTCAGT GAAGATGTCC    2520
450
451   TGCAAGGCTT CTGGCTACAC ATTTACCAGT TACAATATGC ACTGGGTAAA ACAGACACCT   2580
452
453   GGTCGGGGCC TGGAATGGAT TGGAGCTATT TATCCCGGAA ATGGTGATAC TTCCTACAAT   2640
454
455   CAGAAGTTCA AAGGCAAGGC CACATTGACT GCAGACAAAT CCTCCAGCAC AGCCTACATG   2700
456
457   CAGCTCAGCA GCCTGACATC TGAGGACTCT GCGGTCTATT ACTGTGCAAG ATCGACTTAC   2760
458
459   TACGGCGGTG ACTGGTACTT CAATGTCTGG GGCGCAGGGA CCACGGTCAC CGTCTCTGCA   2820
460
461   GCTAGCACCA AGGGCCCATC GGTCTTCCCC CTGGCACCCT CCTCCAAGAG CACCTCTGGG   2880
462
463   GGCACAGCGG CCCTGGGCTG CCTGGTCAAG GACTACTTCC CCGAACCGGT GACGGTGTCG   2940
464
465   TGGAACTCAG GCGCCCTGAC CAGCGGCGTG CACACCTTCC CGGCTGTCCT ACAGTCCTCA   3000
466
467   GGACTCTACT CCCTCAGCAG CGTGGTGACC GTGCCCTCCA GCAGCTTGGG CACCCAGACC   3060
468
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469  TACATCTGCA ACGTGAATCA CAAGCCCAGC AACACCAAGG TGGACAAGAA AGCAGAGCCC  3120
470
471  AAATCTTGTG ACAAAACTCA CACATGCCCA CCGTGCCCAG CACCTGAACT CCTGGGGGGA  3180
472
473  CCGTCAGTCT TCCTCTTCCC CCCAAAACCC AAGGACACCC TCATGATCTC CCGGACCCCT  3240
474
475  GAGGTCACAT GCGTGGTGGT GGACGTGAGC CACGAAGACC CTGAGGTCAA GTTCAACTGG  3300
476
477  TACGTGGACG GCGTGGAGGT GCATAATGCC AAGACAAAGC CGCGGGAGGA GCAGTACAAC  3360
478
479  AGCACGTACC GTGTGGTCAG CGTCCTCACC GTCCTGCACC AGGACTGGCT GAATGGCAAG  3420
480
481  GAGTACAAGT GCAAGGTCTC CAACAAAGCC CTCCCAGCCC CCATCGAGAA AACCATCTCC  3480
482
483  AAAGCCAAAG GGCAGCCCCG AGAACCACAG GTGTACACCC TGCCCCCATC CCGGGATGAG  3540
484
485  CTGACCAAGA ACCAGGTCAG CCTGACCTGC CTGGTCAAAG GCTTCTATCC CAGCGACATC  3600
486
487  GCCGTGGAGT GGGAGAGCAA TGGGCAGCCG GAGAACAACT ACAAGACCAC GCCTCCCGTG  3660
488
489  CTGGACTCCG ACGGCTCCTT CTTCCTCTAC AGCAAGCTCA CCGTGGACAA GAGCAGGTGG  3720
490
491  CAGCAGGGGA ACGTCTTCTC ATGCTCCGTG ATGCATGAGG CTCTGCACAA CCACTACACG  3780
492
493  CAGAAGAGCC TCTCCCTGTC TCCGGGTAAA TGAGGATCCG TTAACGGTTA CCAACTACCT  3840
494
495  AGACTGGATT CGTGACAACA TGCGGCCGTG ATATCTACGT ATGATCAGCC TCGACTGTGC  3900
496
497  CTTCTAGTTG CCAGCCATCT GTTGTTTGCC CCTCCCCCGT GCCTTCCTTG ACCCTGGAAG  3960
498
499  GTGCCACTCC CACTGTCCTT TCCTAATAAA ATGAGGAAAT TGCATCGCAT TGTCTGAGTA  4020
500
501  GGTGTCATTC TATTCTGGGG GGTGGGGTGG GGCAGGACAG CAAGGGGGAG GATTGGGAAG  4080
502
503  ACAATAGCAG GCATGCTGGG GATGCGGTGG GCTCTATGGA ACCAGCTGGG GCTCGACAGC  4140
504
505  GCTGGATCTC CCGATCCCCA GCTTTGCTTC TCAATTTCTT ATTTGCATAA TGAGAAAAAA  4200
506
507  AGGAAAATTA ATTTTAACAC CAATTCAGTA GTTGATTGAG CAAATGCGTT GCCAAAAAGG  4260
508
509  ATGCTTTAGA GACAGTGTTC TCTGCACAGA TAAGGACAAA CATTATTCAG AGGGAGTACC  4320
510
511  CAGAGCTGAG ACTCCTAAGC CAGTGAGTGG CACAGCATTC TAGGGAGAAA TATGCTTGTC  4380
512
513  ATCACCGAAG CCTGATTCCG TAGAGCCACA CCTTGGTAAG GGCCAATCTG CTCACACAGG  4440
514
515  ATAGAGAGGG CAGGAGCCAG GGCAGAGCAT ATAAGGTGAG GTAGGATCAG TTGCTCCTCA  4500
516
517  CATTTGCTTC TGACATAGTT GTGTTGGGAG CTTGGATAGC TTGGACAGCT CAGGGCTGCG  4560
518
519  ATTTCGCGCC AAACTTGACG GCAATCCTAG CGTGAAGGCT GGTAGGATTT TATCCCCGCT  4620
520
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521   GCCATCATGG TTCGACCATT GAACTGCATC GTCGCCGTGT CCCAAAATAT GGGGATTGGC   4680
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523   AAGAACGGAG ACCTACCCTG GCCTCCGCTC AGGAACGAGT TCAAGTACTT CCAAAGAATG   4740
524
525   ACCACAACCT CTTCAGTGGA AGGTAAACAG AATCTGGTGA TTATGGGTAG GAAAACCTGG   4800
526
527   TTCTCCATTC CTGAGAAGAA TCGACCTTTA AAGGACAGAA TTAATATAGT TCTCAGTAGA   4860
528
529   GAACTCAAAG AACCACCACG AGGAGCTCAT TTTCTTGCCA AAAGTTTGGA TGATGCCTTA   4920
530
531   AGACTTATTG AACAACCGGA ATTGGCAAGT AAAGTAGACA TGGTTTGGAT AGTCGGAGGC   4980
532
533   AGTTCTGTTT ACCAGGAAGC CATGAATCAA CCAGGCCACC TTAGACTCTT TGTGACAAGG   5040
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535   ATCATGCAGG AATTTGAAAG TGACACGTTT TTCCCAGAAA TTGATTTGGG GAAATATAAA   5100
536
537   CTTCTCCCAG AATACCCAGG CGTCCTCTCT GAGGTCCAGG AGGAAAAAGG CATCAAGTAT   5160
538
539   AAGTTTGAAG TCTACGAGAA GAAAGACTAA CAGGAAGATG CTTTCAAGTT CTCTGCTCCC   5220
540
541   CTCCTAAAGC TATGCATTTT TATAAGACCA TGGGACTTTT GCTGGCTTTA GATCAGCCTC   5280
542
543   GACTGTGCCT TCTAGTTGCC AGCCATCTGT TGTTTGCCCC TCCCCCGTGC CTTCCTTGAC   5340
544
545   CCTGGAAGGT GCCACTCCCA CTGTCCTTTC CTAATAAAAT GAGGAAATTG CATCGCATTG   5400
546
547   TCTGAGTAGG TGTCATTCTA TTCTGGGGGG TGGGGTGGGG CAGGACAGCA AGGGGGAGGA   5460
548
549   TTGGGAAGAC AATAGCAGGC ATGCTGGGGA TGCGGTGGGC TCTATGGAAC CAGCTGGGGC   5520
550
551   TCGAGCTACT AGCTTTGCTT CTCAATTTCT TATTTGCATA ATGAGAAAAA AAGGAAAATT   5580
552
553   AATTTTAACA CCAATTCAGT AGTTGATTGA GCAAATGCGT TGCCAAAAAG GATGCTTTAG   5640
554
555   AGACAGTGTT CTCTGCACAG ATAAGGACAA ACATTATTCA GAGGGAGTAC CCAGAGCTGA   5700
556
557   GACTCCTAAG CCAGTGAGTG GCACAGCATT CTAGGGAGAA ATATGCTTGT CATCACCGAA   5760
558
559   GCCTGATTCC GTAGAGCCAC ACCTTGGTAA GGGCCAATCT GCTCACACAG GATAGAGAGG   5820
560
561   GCAGGAGCCA GGGCAGAGCA TATAAGGTGA GGTAGGATCA GTTGCTCCTC ACATTTGCTT   5880
562
563   CTGACATAGT TGTGTTGGGA GCTTGGATCG ATCCTCTATG GTTGAACAAG ATGGATTGCA   5940
564
565   CGCAGGTTCT CCGGCCGCTT GGGTGGAGAG GCTATTCGGC TATGACTGGG CACAACAGAC   6000
566
567   AATCGGCTGC TCTGATGCCG CCGTGTTCCG GCTGTCAGCG CAGGGGCGCC CGGTTCTTTT   6060
568
569   TGTCAAGACC GACCTGTCCG GTGCCCTGAA TGAACTGCAG GACGAGGCAG CGCGGCTATC   6120
570
571   GTGGCTGGCC ACGACGGGCG TTCCTTGCGC AGCTGTGCTC GACGTTGTCA CTGAAGCGGG   6180
572
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573    AAGGGACTGG CTGCTATTGG GCGAAGTGCC GGGGCAGGAT CTCCTGTCAT CTCACCTTGC    6240
574
575    TCCTGCCGAG AAAGTATCCA TCATGGCTGA TGCAATGCGG CGGCTGCATA CGCTTGATCC    6300
576
577    GGCTACCTGC CCATTCGACC ACCAAGCGAA ACATCGCATC GAGCGAGCAC GTACTCGGAT    6360
578
579    GGAAGCCGGT CTTGTCGATC AGGATGATCT GGACGAAGAG CATCAGGGGC TCGCGCCAGC    6420
580
581    CGAACTGTTC GCCAGGCTCA AGGCGCGCAT GCCCGACGGC GAGGATCTCG TCGTGACCCA    6480
582
583    TGGCGATGCC TGCTTGCCGA ATATCATGGT GGAAAATGGC CGCTTTTCTG GATTCATCGA    6540
584
585    CTGTGGCCGG CTGGGTGTGG CGGACCGCTA TCAGGACATA GCGTTGGCTA CCCGTGATAT    6600
586
587    TGCTGAAGAG CTTGGCGGCG AATGGGCTGA CCGCTTCCTC GTGCTTTACG GTATCGCCGC    6660
588
589    TCCCGATTCG CAGCGCATCG CCTTCTATCG CCTTCTTGAC GAGTTCTTCT GAGCGGGACT    6720
590
591    CTGGGGTTCG AAATGACCGA CCAAGCGACG CCCAACCTGC CATCACGAGA TTTCGATTCC    6780
592
593    ACCGCCGCCT TCTATGAAAG GTTGGGCTTC GGAATCGTTT TCCGGGACGC CGGCTGGATG    6840
594
595    ATCCTCCAGC GCGGGGATCT CATGCTGGAG TTCTTCGCCC ACCCCAACTT GTTTATTGCA    6900
596
597    GCTTATAATG GTTACAAATA AAGCAATAGC ATCACAAATT TCACAAATAA AGCATTTTTT    6960
598
599    TCACTGCATT CTAGTTGTGG TTTGTCCAAA CTCATCAATC TATCTTATCA TGTCTGGATC    7020
600
601    GCGGCCGCGA TCCCGTCGAG AGCTTGGCGT AATCATGGTC ATAGCTGTTT CCTGTGTGAA    7080
602
603    ATTGTTATCC GCTCACAATT CCACACAACA TACGAGCCGG AAGCATAAAG TGTAAAGCCT    7140
604
605    GGGGTGCCTA ATGAGTGAGC TAACTCACAT TAATTGCGTT GCGCTCACTG CCCGCTTTCC    7200
606
607    AGTCGGGAAA CCTGTCGTGC CAGCTGCATT AATGAATCGG CCAACGCGCG GGGAGAGGCG    7260
608
609    GTTTGCGTAT TGGGCGCTCT TCCGCTTCCT CGCTCACTGA CTCGCTGCGC TCGGTCGTTC    7320
610
611    GGCTGCGGCG AGCGGTATCA GCTCACTCAA AGGCGGTAAT ACGGTTATCC ACAGAATCAG    7380
612
613    GGGATAACGC AGGAAAGAAC ATGTGAGCAA AAGGCCAGCA AAAGGCCAGG AACCGTAAAA    7440
614
615    AGGCCGCGTT GCTGGCGTTT TTCCATAGGC TCCGCCCCCC TGACGAGCAT CACAAAAATC    7500
616
617    GACGCTCAAG TCAGAGGTGG CGAAACCCGA CAGGACTATA AAGATACCAG GCGTTTCCCC    7560
618
619    CTGGAAGCTC CCTCGTGCGC TCTCCTGTTC CGACCCTGCC GCTTACCGGA TACCTGTCCG    7620
620
621    CCTTTCTCCC TTCGGGAAGC GTGGCGCTTT CTCAATGCTC ACGCTGTAGG TATCTCAGTT    7680
622
623    CGGTGTAGGT CGTTCGCTCC AAGCTGGGCT GTGTGCACGA ACCCCCCGTT CAGCCCGACC    7740
624
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625   GCTGCGCCTT ATCCGGTAAC TATCGTCTTG AGTCCAACCC GGTAAGACAC GACTTATCGC   7800
626
627   CACTGGCAGC AGCCACTGGT AACAGGATTA GCAGAGCGAG GTATGTAGGC GGTGCTACAG   7860
628
629   AGTTCTTGAA GTGGTGGCCT AACTACGGCT ACACTAGAAG GACAGTATTT GGTATCTGCG   7920
630
631   CTCTGCTGAA GCCAGTTACC TTCGGAAAAA GAGTTGGTAG CTCTTGATCC GGCAAACAAA   7980
632
633   CCACCGCTGG TAGCGGTGGT TTTTTTGTTT GCAAGCAGCA GATTACGCGC AGAAAAAAAG   8040
634
635   GATCTCAAGA AGATCCTTTG ATCTTTTCTA CGGGGTCTGA CGCTCAGTGG AACGAAAACT   8100
636
637   CACGTTAAGG GATTTTGGTC ATGAGATTAT CAAAAAGGAT CTTCACCTAG ATCCTTTTAA   8160
638
639   ATTAAAAATG AAGTTTTAAA TCAATCTAAA GTATATATGA GTAAACTTGG TCTGACAGTT   8220
640
641   ACCAATGCTT AATCAGTGAG GCACCTATCT CAGCGATCTG TCTATTTCGT TCATCCATAG   8280
642
643   TTGCCTGACT CCCCGTCGTG TAGATAACTA CGATACGGGA GGGCTTACCA TCTGGCCCCA   8340
644
645   GTGCTGCAAT GATACCGCGA GACCCACGCT CACCGGCTCC AGATTTATCA GCAATAAACC   8400
646
647   AGCCAGCCGG AAGGGCCGAG CGCAGAAGTG GTCCTGCAAC TTTATCCGCC TCCATCCAGT   8460
648
649   CTATTAATTG TTGCCGGGAA GCTAGAGTAA GTAGTTCGCC AGTTAATAGT TTGCGCAACG   8520
650
651   TTGTTGCCAT TGCTACAGGC ATCGTGGTGT CACGCTCGTC GTTTGGTATG GCTTCATTCA   8580
652
653   GCTCCGGTTC CCAACGATCA AGGCGAGTTA CATGATCCCC CATGTTGTGC AAAAAAGCGG   8640
654
655   TTAGCTCCTT CGGTCCTCCG ATCGTTGTCA GAAGTAAGTT GGCCGCAGTG TTATCACTCA   8700
656
657   TGGTTATGGC AGCACTGCAT AATTCTCTTA CTGTCATGCC ATCCGTAAGA TGCTTTTCTG   8760
658
659   TGACTGGTGA GTACTCAACC AAGTCATTCT GAGAATAGTG TATGCGGCGA CCGAGTTGCT   8820
660
661   CTTGCCCGGC GTCAATACGG GATAATACCG CGCCACATAG CAGAACTTTA AAAGTGCTCA   8880
662
663   TCATTGGAAA ACGTTCTTCG GGGCGAAAAC TCTCAAGGAT CTTACCGCTG TTGAGATCCA   8940
664
665   GTTCGATGTA ACCCACTCGT GCACCCAACT GATCTTCAGC ATCTTTTACT TTCACCAGCG   9000
666
667   TTTCTGGGTG AGCAAAAACA GGAAGGCAAA ATGCCGCAAA AAAGGGAATA AGGGCGACAC   9060
668
669   GGAAATGTTG AATACTCATA CTCTTCCTTT TTCAATATTA TTGAAGCATT TATCAGGGTT   9120
670
671   ATTGTCTCAT GAGCGGATAC ATATTTGAAT GTATTTAGAA AAATAAACAA ATAGGGGTTC   9180
672
673   CGCGCACATT TCCCCGAAAA GTGCCACCT   9209
674
675
676   (4)INFORMATION FOR SEQ ID NO: 3:

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677
678    (i)SEQUENCE CHARACTERISTICS:
679
680    (A)LENGTH:  47 bases
681    (B)TYPE:  nucleic acid
682    (C)STRANDEDNESS:  single
683    (D)TOPOLOGY:  linear
684
685    (ii)MOLECULE TYPE:  DNA (genomic)
686
687    (iii)HYPOTHETICAL:  no
688
689    (iv)ANTI-SENSE:  no
690
691    (ix)SEQUENCE DESCRIPTION:  SEQ ID NO:  3:
692
693    ATC ACA GAT CTC TCA CCA TGG ATT TTC AGG TBC AGA TTA TCA GCT45
694    TC47
695    (5) INFORMATION FOR SEQ ID NO:  4:
696
697    (i)SEQUENCE CHARACTERISTICS:
698
699    (A)LENGTH:  30 bases
700    (B)TYPE:  nucleic acid
701    (C)STRANDEDNESS:  single
702    (D)TOPOLOGY:  linear
703
704    (ii)MOLECULE TYPE:  DNA (genomic)
705
706    (iii)HYPOTHETICAL:  no
707
708    (iv)ANTI-SENSE:  yes
709
710    (ix)SEQUENCE DESCRIPTION:  SEQ ID NO:  4:
711
712    TGC AGC ATC CGT ACG TTT GAT TTC CAG CTT 30
713
714    (6)INFORMATION FOR SEQ ID NO:  5:
715
716    (i)SEQUENCE CHARACTERISTICS:
717
718    (A)LENGTH:  384 bases
719    (B)TYPE:  nucleic acid
720    (C)STRANDEDNESS:  single
721    (D)TOPOLOGY:  linear
722
723    (ii)MOLECULE TYPE:  DNA (genomic)
724
725    (iii)HYPOTHETICAL:  no
726
727    (iv)ANTI-SENSE:  no
728
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[Handwritten annotation: "Space before the nucleic count number"]

[Handwritten annotation near line 696: "Space"]

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729   (ix)SEQUENCE DESCRIPTION:   SEQ ID NO:   5:
730
731
732     ATG GAT TTT CAG GTG CAG ATT ATC AGC TTC CTG CTA ATC AGT GCT TCA GTC 51
733
734     ATA ATG TCC AGA GGG CAA ATT GTT CTC TCC CAG TCT CCA GCA ATC CTG TCT102
735
736     GCA TCT CCA GGG GAG AAG GTC ACA ATG ACT TGC AGG GCC AGC TCA AGT GTA153
737
738     AGT TAC ATC CAC TGG TTC CAG CAG AAG CCA GGA TCC TCC CCC AAA CCC TGG204
739
740     ATT TAT GCC ACA TCC AAC CTG GCT TCT GGA GTC CCT GTT CGC TTC AGT GGC255
741
742     AGT GGG TCT GGG ACT TCT TAC TCT CTC ACA ATC AGC AGA GTG GAG GCT GAA306
743
744     GAT GCT GCC ACT TAT TAC TGC CAG CAG TGG ACT AGT AAC CCA CCC ACG TTC357
745
746     GGA GGG GGG ACC AAG CTG GAA ATC AAA384
747
748
749   (7)INFORMATION FOR SEQ ID NO:   6:
750
751   (i)SEQUENCE CHARACTERISTICS:
752
753   (A)LENGTH:   27 bases
754   (B)TYPE:   nucleic acid
755   (C)STRANDEDNESS:   single
756   (D)TOPOLOGY:   linear
757
758   (ii)MOLECULE TYPE:   DNA (genomic)
759
760   (iii)HYPOTHETICAL:   no
761
762   (iv)ANTI-SENSE:   no
763
764   (ix)SEQUENCE DESCRIPTION:   SEQ ID NO:   6:
765
766   GCG GCT CCC ACG CGT GTC CTG TCC CAG27
767
768   (8)INFORMATION FOR SEQ ID NO:   7:
769
770   (i)SEQUENCE CHARACTERISTICS:
771
772   (A)LENGTH:   29 bases
773   (B)TYPE:   nucleic acid
774   (C)STRANDEDNESS:   single
775   (D)TOPOLOGY:   linear
776
777   (ii)MOLECULE TYPE:   DNA (genomic)
778
779   (iii)HYPOTHETICAL:   no
780
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781  (iv)ANTI-SENSE:  yes
782
783  (ix)SEQUENCE DESCRIPTION:   SEQ ID NO:  7:
784
785  GGS TGT TGT GCT AGC TGM RGA GAC RGT GA29
786
787  (9)INFORMATION FOR SEQ ID NO:  8:
788
789  (i)SEQUENCE CHARACTERISTICS:
790
791  (A)LENGTH:   420 bases
792  (B)TYPE:   nucleic acid
793  (C)STRANDEDNESS:   single
794  (D)TOPOLOGY:   linear
795
796  (ii)MOLECULE TYPE:   DNA (genomic)
797
798  (iii)HYPOTHETICAL:   no
799
800  (iv)ANTI-SENSE:   no
801
802  (ix)SEQUENCE DESCRIPTION:   SEQ ID NO:  8:
803
804  ATG GGT TGG AGC CTC ATC TTG CTC TTC CTT GTC GCT GTT GCT ACG CGT GTC 51
805
806  CTG TCC CAG GTA CAA CTG CAG CAG CCT GGG GCT GAG CTG GTG AAG CCT GGG102
807
808  GCC TCA GTG AAG ATG TCC TGC AAG GCT TCT GGC TAC ACA TTT ACC AGT TAC153
809
810  AAT ATG CAC TGG GTA AAA CAG ACA CCT GGT CGG GGC CTG GAA TGG ATT GGA 204
811
812  GCT ATT TAT CCC GGA AAT GGT GAT ACT TCC TAC AAT CAG AAG TTC AAA GGC255
813
814  AAG GCC ACA TTG ACT GCA GAC AAA TCC TCC AGC ACA GCC TAC ATG CAG CTC306
815
816  AGC AGC CTG ACA TCT GAG GAC TCT GCG GTC TAT TAC TGT GCA AGA TCG ACT357
817
818  TAC TAC GGC GGT GAC TGG TAC TTC AAT GTC TGG GGC GCA GGG ACC ACG GTC408
819  ACC GTC TCT GCA        420
820
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